SEQUENCE LISTING

(1) GENERAL INFORMATION:

    (i) APPLICANT:
        (A) NAME: Geron Corporation
        (B) STREET: 230 Constitution Drive
        (C) CITY: Menlo Park
        (D) STATE: California
        (E) POSTAL CODE (ZIP): 94025
        (F) TELEPHONE: (415) 473-7700
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        (H) TELEX:

    (i) APPLICANT:
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        (B) STREET: 541 Winterberry Way
        (C) CITY: San Jose
        (D) STATE: California
        (E) POSTAL CODE (ZIP): 95129
        (F) TELEPHCNE:
        (G) TELEFAX:
        (H) TELEX:

    (i) APPLICANT:
        (A) NAME: Wu, Fred
        (B) STREET: 1143 Greene Terrace
        (C) CITY: Davis
        (D) STATE: California
        (E) POSTAL CODE (ZIP): 94616
        (F) TELEPHONE:
        (G) TELEFAX:
        (H) TELEX:

    (i) APPLICANT:
        (A) NAME: Kealey, James T.
        (B) STREET: 9 Mono Lane
        (C) CITY: San Anselmo
        (D) STATE: California
        (E) POSTAL CODE (ZIP): 94960
        (F) TELEPHONE:
        (G) TELEFAX:
        (H) TELEX:

    (i) APPLICANT:
        (A) NAME: Pruzan, Ronald
        (B) STREET: 852 Matadero Avenue
        (C) CITY: Palo Alto
        (D) STATE: California
        (E) POSTAL CODE (ZIP): 94306
        (F) TELEPHONE:
        (G) TELEFAX:
        (H) TELEX:

    (i) APPLICANT:
        (A) NAME: Weinrich, Scott L.

(B) STREET: 1830 Brewster Avenue
           (C) CITY: Redwood City
           (D) STATE: California
           (E) POSTAL CODE (ZIP): 94062
           (F) TELEPHONE:
           (G) TELEFAX:
           (H) TELEX:

     (ii) TITLE OF INVENTION: Methods for Detecting and Inhibiting the
           RNA Component of Telomerase

    (iii) NUMBER OF SEQUENCES: 29

     (iv) CORRESPONDENCE ADDRESS:
           (A) ADDRESSEE: Townsend and Townsend and Crew LLP
           (B) STREET: Two Embarcadero Center, Eighth Floor
           (C) CITY: San Francisco
           (D) STATE: California
           (E) COUNTRY: USA
           (F) ZIP: 94111-3834

      (v) COMPUTER READABLE FORM:
           (A) MEDIUM TYPE: Floppy disk
           (B) COMPUTER: IBM PC compatible
           (C) OPERATING SYSTEM: PC-DOS/MS-DOS
           (D) SOFTWARE: PatentIn Release #1.0, Version #1.30

     (vi) CURRENT APPLICATION DATA:
           (A) APPLICATION NUMBER: WO PCT/US97/Not yet assigned
           (B) FILING DATE: Not yet assigned
           (C) CLASSIFICATION:

    (vii) PRIOR APPLICATION DATA:
           (A) APPLICATION NUMBER: US 08/770,564
           (B) FILING DATE: 20-DEC-1996

    (vii) PRIOR APPLICATION DATA:
           (A) APPLICATION NUMBER: US 08/770,565
           (B) FILING DATE: 20-DEC-1996

   (viii) ATTORNEY/AGENT INFORMATION:
           (A) NAME: Storella, John R.
           (B) REGISTRATION NUMBER: 32,944
           (C) REFERENCE/DOCKET NUMBER: 15389-27PC

     (ix) TELECOMMUNICATION INFORMATION:
           (A) TELEPHONE: (415) 576-0200
           (B) TELEFAX: (415) 576-0300

(2) INFORMATION FOR SEQ ID NO:1:

      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 981 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear

.(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:
    (A) NAME/KEY: -
    (B) LOCATION: 1..981
    (D) OTHER INFORMATION: /note= "PstI fragment of
                              SauIIIA1-HindIII fragment of clone 28-1
                              containing the sequence of the RNA
                              component of human telomerase (hTR)"

(ix) FEATURE:
    (A) NAME/KEY: misc_RNA
    (B) LOCATION: 266..716
    (D) OTHER INFORMATION: /product= "RNA component of human
                              telomerase (hTR)"

(ix) FEATURE:
    (A) NAME/KEY: misc_feature
    (B) LOCATION: 311..320
    (D) OTHER INFORMATION: /note= "template region of hTR"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

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CTGCAGAGGA TAGAAAAAAG GCCCTCTGAT ACCTCAAGTT AGTTTCACCT TTAAAGAAGG        60

TCGGAAGTAA AGACGCAAAG CCTTTCCCGG ACGTGCGGAA GGGCAACGTC CTTCCTCATG       120

GCCGGAAATG GAACTTTAAT TTCCCGTTCC CCCCAACCAG CCCGCCCGAG AGAGTGACTC       180

TCACGAGAGC CGCGAGAGTC AGCTTGGCCA ATCCGTGCGG TCGGCGGCCG CTCCCTTTAT       240

AAGCCGACTC GCCCGGCAGC GCACCGGGTT GCGGAGGGTG GGCCTGGGAG GGGTGGTGGC       300

CATTTTTTGT CTAACCCTAA CTGAGAAGGG CGTAGGCGCC GTGCTTTTGC TCCCCGCGCG       360

CTGTTTTTCT CGCTGACTTT CAGCGGGCGG AAAAGCCTCG GCCTGCCGCC TTCCACCGTT       420

CATTCTAGAG CAAACAAAAA ATGTCAGCTG CTGGCCCGTT CGCCCCTCCC GGGGACCTGC       480

GGCGGGTCGC CTGCCCAGCC CCCGAACCCC GCCTGGAGGC CGCGGTCGGC CCGGGGCTTC       540

TCCGGAGGCA CCCACTGCCA CCGCGAAGAG TTGGGCTCTG TCAGCCGCGG GTCTCTCGGG       600

GGCGAGGGCG AGGTTCAGGC CTTTCAGGCC GCAGGAAGAG GAACGGAGCG AGTCCCCGCG       660

CGCGGCGCGA TTCCCTGAGC TGTGGGACGT GCACCCAGGA CTCGGCTCAC ACATGCAGTT       720

CGCTTTCCTG TTGGTGGGGG GAACGCCGAT CGTGCGCATC CGTCACCCCT CGCCGGCAGT       780

GGGGGCTTGT GAACCCCCAA ACCTGACTGA CTGGGCCAGT GTGCTGCAAA TTGGCAGGAG       840

ACGTGAAGGC ACCTCCAAAG TCGGCCAAAA TGAATGGGCA GTGAGCCGGG GTTGCCTGGA       900

GCCGTTCCTG CGTGGGTTCT CCCGTCTTCC GCTTTTTGTT GCCTTTTATG GTTGTATTAC       960
```

AACTTAGTTC CTGCTCTGCA G   981

(2) INFORMATION FOR SEQ ID NO:2:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 30 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (ii) MOLECULE TYPE: DNA

        (ix) FEATURE:
            (A) NAME/KEY: -
            (B) LOCATION: 1..30
            (D) OTHER INFORMATION: /note= "oligo 14"

        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

CGTTCCTCTT CCTGCGGCCT GAAACGGTGA   30

(2) INFORMATION FOR SEQ ID NO:3:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 20 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (ii) MOLECULE TYPE: DNA

        (ix) FEATURE:
            (A) NAME/KEY: -
            (B) LOCATION: 1..20
            (D) OTHER INFORMATION: /note= "oligo 14ab"

        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

CGTTCCTCTT CCTGCGGCCT   20

(2) INFORMATION FOR SEQ ID NO:4:

        (i) SEQUENCE CHARACTERISTICS:
            (A) LENGTH: 12 base pairs
            (B) TYPE: nucleic acid
            (C) STRANDEDNESS: single
            (D) TOPOLOGY: linear

        (ii) MOLECULE TYPE: DNA

```
(ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..12
        (D) OTHER INFORMATION: /note= "oligo 14d"


(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CGTTCCTCTT CC                                                                    12


(2) INFORMATION FOR SEQ ID NO:5:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 30 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..30
        (D) OTHER INFORMATION: /note= "oligo 16"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

CTGACAGAGC CCAACTCTTC GCGGTGGCAG                                                 30


(2) INFORMATION FOR SEQ ID NO:6:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /note= "oligo 16ab"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

CTGACAGAGC CCAACTCTTC                                                            20
```

(2) INFORMATION FOR SEQ ID NO:7:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: DNA

   (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /note= "oligo 16bc"

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

CCAACTCTTC GCGGTGGCAG   20

(2) INFORMATION FOR SEQ ID NO:8:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 30 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: DNA

   (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..30
        (D) OTHER INFORMATION: /note= "oligo 21"

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

GCTCTAGAAT GAACGGTGGA AGGCGGCAGG   30

(2) INFORMATION FOR SEQ ID NO:9:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 19 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: DNA

   (ix) FEATURE:
        (A) NAME/KEY: -

(B) LOCATION: 1..19
          (D) OTHER INFORMATION: /note= "oligo 21ab"

     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

GCTCTAGAAT GAACGGTGG   19

(2) INFORMATION FOR SEQ ID NO:10:

     (i). SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 15 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

     (ii) MOLECULE TYPE: DNA

     (ix) FEATURE:
          (A) NAME/KEY: -
          (B) LOCATION: 1..15
          (D) OTHER INFORMATION: /note= "oligo 21ab3"

     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

GCTCTAGAAT GAACG   15

(2) INFORMATION FOR SEQ ID NO:11:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 11 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

     (ii) MOLECULE TYPE: DNA

     (ix) FEATURE:
          (A) NAME/KEY: -
          (B) LOCATION: 1..11
          (D) OTHER INFORMATION: /note= "oligo 21ab2"

     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

GCTCTAGAAT G   11

(2) INFORMATION FOR SEQ ID NO:12:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 7 base pairs

```
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..7
        (D) OTHER INFORMATION: /note= "oligo 21ab1"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

GCTCTAG                                                          7


(2) INFORMATION FOR SEQ ID NO:13:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..20
        (D) ·OTHER INFORMATION: /note= "oligo 20/21"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

CATTTTTTGT TTGCTCTAGA                                            20


(2) INFORMATION FOR SEQ ID NO:14:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 17 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..17
        (D) OTHER INFORMATION: /note= "polynucleotide RP2"
```

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

CGGGCCAGCA GCTGACA   17

(2) INFORMATION FOR SEQ ID NO:15:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 26 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

     (ii) MOLECULE TYPE: DNA

     (ix) FEATURE:
          (A) NAME/KEY: -
          (B) LOCATION: 1..26
          (D) OTHER INFORMATION: /note= "28S rRNA forward primer"

     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

GCTAAATACC GGCACGAGAC CGATAG   26

(2) INFORMATION FOR SEQ ID NO:16:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 20 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

     (ii) MOLECULE TYPE: DNA

     (ix) FEATURE:
          (A) NAME/KEY: -
          (B) LOCATION: 1..20
          (D) OTHER INFORMATION: /note= "28S rRNA forward primer"

     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

GCTAAATACC GGCACGAGAC   20

(2) INFORMATION FOR SEQ ID NO:17:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 18 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

(2) INFORMATION FOR SEQ ID NO:20:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 26 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA

    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..26
        (D) OTHER INFORMATION: /note= "28S rRNA reverse primer"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

GGTTTCACGC CCTCTTGAAC TCTCTC   26

(2) INFORMATION FOR SEQ ID NO:21:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 21 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA

    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..21
        (D) OTHER INFORMATION: /note= "28S rRNA reverse primer"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

GTTTCACGCC CTCTTGAACT C   21

(2) INFORMATION FOR SEQ ID NO:22:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 19 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA

    (ix) FEATURE:

```
        (A) NAME/KEY: -
        (B) LOCATION: 1..19
        (D) OTHER INFORMATION: /note= "28S rRNA reverse primer"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

GTTTCACGCC CTCTTGAAC                                              19


(2) INFORMATION FOR SEQ ID NO:23:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 19 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..19
        (D) OTHER INFORMATION: /note= "28S rRNA reverse primer"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

CTAGTTTCAC GCCCTCTTG                                              19


(2) INFORMATION FOR SEQ ID NO:24:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 19 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA


    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..19
        (D) OTHER INFORMATION: /note= "28S rRNA reverse primer"


    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

CTGGTTTCAC GCCCTCTTG                                              19


(2) INFORMATION FOR SEQ ID NO:25:

    (i) SEQUENCE CHARACTERISTICS:
```

(A) LENGTH: 26 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA

    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..26
        (D) OTHER INFORMATION: /note= "hTR forward primer"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

GAAGGGCGTA GGCGCCGTGC TTTTGC   26

(2) INFORMATION FOR SEQ ID NO:26:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 27 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA

    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..27
        (D) OTHER INFORMATION: /note= "hTR reverse primer"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

GTTTGCTCTA GAATGAACGG TGGAAGG   27

(2) INFORMATION FOR SEQ ID NO:27:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 base pairs
        (B) TYPE: nucleic acid
        (C) STRANDEDNESS: single
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: DNA

    (ix) FEATURE:
        (A) NAME/KEY: -
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /note= "oligo 14bc"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

CCTGCGGCCT GAAACGGTGA   20

(2) INFORMATION FOR SEQ ID NO:28:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 10 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

     (ii) MOLECULE TYPE: DNA (genomic)

     (ix) FEATURE:
          (A) NAME/KEY: -
          (B) LOCATION: 1..10
          (D) OTHER INFORMATION: /note= "template region of hTR"

     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

CTAACCCTAA   10

(2) INFORMATION FOR SEQ ID NO:29:

     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 451 base pairs
          (B) TYPE: nucleic acid
          (C) STRANDEDNESS: single
          (D) TOPOLOGY: linear

     (ii) MOLECULE TYPE: DNA (genomic)

     (ix) FEATURE:
          (A) NAME/KEY: -
          (B) LOCATION: 1..451
          (D) OTHER INFORMATION: /note= "RNA component of human
                              telomerase (hTR)"

     (ix) FEATURE:
          (A) NAME/KEY: misc_feature
          (B) LOCATION: 46..55
          (D) OTHER INFORMATION: /note= "template region of hTR"

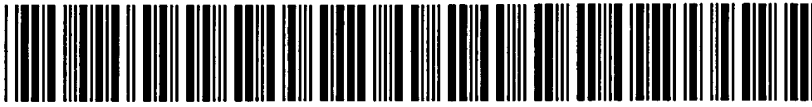   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

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GGCGGAAAAG CCTCGGCCTG CCGCCTTCCA CCGTTCATTC TAGAGCAAAC AAAAAATGTC        180

AGCTGCTGGC CCGTTCGCCC CTCCCGGGGA CCTGCGGCGG GTCGCCTGCC CAGCCCCCGA        240

ACCCCGCCTG GAGGCCGCGG TCGGCCCGGG GCTTCTCCGG AGGCACCCAC TGCCACCGCG        300

AAGAGTTGGG CTCTGTCAGC CGCGGGTCTC TCGGGGGCGA GGGCGAGGTT CAGGCCTTTC        360

AGGCCGCAGG AAGAGGAACG GAGCGAGTCC CCGCGCGCGG CGCGATTCCC TGAGCTGTGG        420

GACGTGCACC CAGGACTCGG CTCACACATG C                                      451
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